



1600

RAW SEQUENCE LISTING DATE: 04/11/2003 FATENT APPLICATION: US/09/816,697A TIME: 14:30:02

Imput Set : A:\EP.txt

Output Set: N:\CRF4\04112003\I816697A.raw

```
5 K110 APPLICANT: Lorenz, M.
     9 - 120 - TITLE OF INVENTION: A NOVEL P-SELECTIN GLYCOPROTEIN LIGAND (PSGL-1) BINDING
PROTEIN AND USES
              THEREFOR.
     14 <130 / FILE FEFERENCE: 08702.0005-00000
     15 -: 140 - CURRENT APPLICATION NUMBER: 09/816,697A
     PG -: 141 - CURRENT FILING DATE: 2001-03-23
     14 <150 - PRIOR APPLICATION NUMBER: 60/192,104
     16 0151 / PRIOR FILING DATE: 2000-03-24
     30 -1160 - NUMBER OF SEQ ID NOS: 6
                                                               ENTERED
     34 K170 - SOFTWARE: PatentIn version 3.1
     38 KO10 - SEQ ID NO: 1
     40 -211 - LENGTH: 951
     42 KI 10× TYPE: DNA
     44 CL13 · OFGANISM: Homo sabiers
     48 K400> SEQUENCE: 1
                                                                                50
     49 atggsaagte sagagsacee tyggagseet ggetgsatgg gacssataac ceagtgsacg
                                                                                100
     51 graaggadoo agraaggaago accagdosat ggoocogado toocgcacco aggacotgad:
     83 gugdaettag udabadadag tiggdetgago tedaactoda geatgaddac gegiggagett.
                                                                                340
     55 cagcagtact ggcagaacca gaaatgoogo tggaagcacg tcaaactgot ctttgagato
                                                                                200
     57 gottoagoto goatogagga gagaaaagto totaagttttg tggtgtaboa aatoatogto.
     3.9 atomaganty ggagetitya maamaamaay googtootgy aanggogeta ttomgantto
                                                                                360
                                                                                400
     ol gogaagetee agaaageget getgaagaeg tteagggagg agategaaga egtggagttt.
                                                                               40
     63 occaggaage achtgaetgg gaaetteget gaggagatga tetgtgageg teggegegee
                                                                                540
     W engaggagt weetiggyeet getelacged atcogetige tigggeegete coggraphic
     67 otgqwettee temegoggee ggagetgege gaggettteg getgeetgeg ggeeggeeag
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     na twocogogog contiguaget gethotigogo gitgoligoego igoaggagaa geloacegoo
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     71 eactgoesty egocogologic energy eacty typicoty typitytycca cogogolote
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     7% maconoccey negaggeott oyoggoogga gagagggood tgcagegoot gcaggoodgg
     7% gagggocate getactatge geetetgetg gabgocatgg teogcotggs etabgogotg
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     77 ggcaaggact togtquotot gcaqqagagg otggaggaga gocaqotoog gaggoodaog
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     79 opporagges toscoptigas gyagoticset gtgogagast scotgosotig a
    BU KUTON SEQ ID NO: 2
    84 MAIL DENGTH: 316
     86 - 212: TYPE: FET
    88 HILLS: OFGANISM: Home sapiers
     9. 4400 SEQUENCE: 1
     94 Met Ala Ser Pro Glu His Pro Gly Ser Pro Gly Cys Met Gly Pro Ile
     95 1
                                             10
     96 Thr Gln Cys Thr Ala Arg Thr Gln Gln Glu Ala Pro Ala Thr Gly Pro
     1.1
     102 Asp Leu Fro His Pro Gly Fro Asp Gly His Leu Asp Thr His Ser Gly
                                      40
```

106 Leu Ser Ser Asn Ser Ser Met Thr Thr Arg Glu Leu Gln Gln Tyr Trp

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107		50					55					60					
	Gln	Asn	Gln	1.78	Cys	Arg	$T^{i}r^{i}p^{i}$	Lys	His	Val	Lys	Leu	Leu	Ph∈	Glu	Ile	
111						70.					7.5					80	
114	Ala	Ser	Ala	Arq		Glu	Blu	Arg	Lуз		S⊕r	Lys	Phe	Val		Tyr	
11:					35					90					95		
	Gln	lle	1.1e		11⊕	Gln	Thr	GLY		Phe	Азр	Asn	Aar.		Ala	Va.I	
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	Lesa	Glu		Arq	Tyr	Ser	ASP		ALâ	-уз	Leu	G.l.n		Ala	Leu	7.69.7	
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	S_{t+1}	Δνα	(21s)	Phe		Aeri	Dh e	[145]	Thir		Pro	Glu	Len	Ara		Ala	
139		117.7	Tallah ak	180	шчи	, 7.º F.		٠,٠	185	**** =		J 3. G	21.5.3	190	54.5		
	Phe	Glv	rivs		Ara	Ala	Glv	Gln		Pro	Ara	Ala	Leu		Leu	Leu	
14.		y	195		3			200	- 1 -				205				
	Leu	Ara		โดน	Pro	Leu	Glr.	Glu	Lys	Leu	Tr.r	Ala	Eis	Cys	Pro	Ala	
147		210					.15		•			220		-			
150	Ala	Ala	Val	Pino	Ala	Leu	Cys	Ala	Val	Leu	Leu	Суз	Eis	Arg	Asp	Leu	
151	25					230					235					240	
154	Asp	Arg	Pro	Alla	Glu	Ala	Phe	Ala	Ala	Gly	Glu	Arg	Ala	Leu	Gl.r.	Arg	
155					245					250					255		
	Lou	Glr.	Ala		Glu	G1Y	His	Arg		Tyr	Ala	Pro	L∈u		Asp	Ala	
159				260			_		265		_			270	_	- 1	
	Met	Vāl.		Leu	Ala	Tyr	Ala		GLy	Lys	Asp	Phie		Thr	ьeu	GIn	
163			175	~ 1	2.1		~ 1	280	70	70 .	Б	m1	285	7)		T1 ~	
	Glu	-	≟⊖u	Glu	Blu	ser		Leu	Arg	Arg	Pro	300	Pro	Arg	וΣ⊤Σ	116	
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		305 310 315 310 315															
		KOIO - SEQ ID NO: 3 KLII- BENGTH: 34															
		Q:12 - TYPE: DNA Q:13 - ORGANISM: Artificial sequence															
		1230 - FEATURE:															
		UN3 - OTHER INFORMATION: Frimer															
		4400 - SEQUENCE: 3															
	atac					aa aa	gcaac	gggoo	c aca	at							34
						-											
	<.011																
	SHONESH TYPE: DNA																
					Arti	Lfici	lal s	seque	ence								
201	+213 + ORGANISM: Artificial sequence +220 + FEATURE:																
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	-(400)																
207	at.ac	cagga	atc o	cagaç	gtgaç	ge ta	aggg	gagga	a aag	3							33

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210 <210 > SEQ ID NO: 5 313 4311 LENGTH: 15 D14 HH12 TYPE: PRT [16] M.:13 · ORGANISM: Artificial sequence .20 -1.20 - FEATURE: LANGE OTHER INFORMATION: antigen .114 -1400 - SEQUENCE: 5 000 Gln Glu Arg Leu Glu Glu Ser Gln Leu Arg Arg Pro Thr Pro Arg 227 1 E. 10 350 -110 - SEQ 1D NO: 6 232 - 211 - DENGTH: 11 334 + 3212 + TYPE: PFT236 -013 - ORGANISM: Artificial sequence 240 KIRCO FEATURE: 140 -0023 OTHER INFORMATION: T7 tag 244 - 400 > SEQUENCE: 6 246 Met Ala Ser Met Thr Gly Gly Gln Gln Met Gly 247 1

RAW SEQUENCE LISTING ERROR SUMMARY
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Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 9

VERIFICATION SUMMARY

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